

## Product datasheet for RC212367L4V

## OriGene Technologies, Inc.

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## CYP3A43 (NM\_057096) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** CYP3A43 (NM\_057096) Human Tagged ORF Clone Lentiviral Particle

**Symbol:** CYP3A43

Mammalian Cell Puromycin

Selection:

**Vector:** pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

**ACCN:** NM\_057096

**ORF Size:** 1260 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC212367).

OTI Disclaimer:

Sequence:

The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. More info

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

**RefSeq:** <u>NM 057096.2</u>

 RefSeq Size:
 2001 bp

 RefSeq ORF:
 1263 bp

 Locus ID:
 64816

 UniProt ID:
 Q9HB55

 Cytogenetics:
 7q22.1

**Protein Families:** Druggable Genome, Transmembrane

**Protein Pathways:** Drug metabolism - cytochrome P450, Drug metabolism - other enzymes, Linoleic acid

metabolism, Metabolic pathways, Metabolism of xenobiotics by cytochrome P450, Retinol

metabolism





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MW: 48.1 kDa

**Gene Summary:** This gene encodes a member of the cytochrome P450 superfamily of enzymes. The

cytochrome P450 proteins are monooxygenases which catalyze many reactions involved in drug metabolism and synthesis of cholesterol, steroids and other lipids. The encoded protein has a low level of testosterone hydroxylase activity, and may play a role in aging mechanisms

and cancer progression. This gene is part of a cluster of cytochrome P450 genes on chromosome 7q21.1. Alternate splicing results in multiple transcript variants encoding

different isoforms. [provided by RefSeq, Jul 2013]